

II. AMENDMENTS TO THE CLAIMS

1-15. (Canceled)

16. (New) A method for production of a L-amino acid derived from a beta-aryl-substituted L-amino acid, comprising:

(a) fermenting an *E.coli* host cell that contains an isolated polynucleotide selected from the group consisting of

(i) a nucleotide sequence as set forth in SEQ ID NO: 1; and

(ii) a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2;

(b) expressing an *Arthrobacter aurescens*' L-N-carbamoylase from step (a); and

(c) contacting the L-N carbamoylase of step (b) with N-carbamoyl or N-formyl amino acids to produce said L-amino acid derived from a beta-aryl-substituted L-amino acid.

17. (New) The method according to claim 16, further comprising the step of immobilizing the L-N-carbamoylase onto carriers.

18. (New) The method according to claim 17, wherein the L-N-carbamoylase is covalently immobilized on EAH-sepharose.

19. (New) The method according to claim 16, wherein the induction of expression of L-N-carbamoylase is by rhamnose, IPTG, or lactose.

20. (New) The method according to claim 16, wherein N-formyl-D,L tryptophase, N-acetyl-D,L-tryptophane, and N-carbamoyl-D, L-phenylalanine serve as substrates for the L-N-carbamoylase.

21. (New) The method according to claim 16, wherein the isolated polynucleotide is the *hyuC* gene of *Arthrobacter aurescens*.

22. (New) A method for production of L-methionine comprising:

(a) fermenting an *E.coli* host cell that contains an isolated polynucleotide selected from the group consisting of

- (i) a nucleotide sequence as set forth in SEQ ID NO: 1; and
- (ii) a nucleotide sequence encoding the polypeptide as set forth in SEQ ID NO: 2;
- (b) producing an *Arthrobacter aurescens*' L-N-carbamoylase from step (a); and
- (c) contacting the L-N carbamoylase of step (b) with N-carbamoyl-L-thienylalanine to produce L-methionine.

23. (New) The method according to claim 22, further comprising the step of immobilizing the L-N-carbamoylase onto carriers.

24. (New) The method according to claim 23, wherein the L-N-carbamoylase is covalently immobilized on EAH-sepharose.

25. (New) The method according to claim 22, wherein the induction of expression of L-N-carbamoylase is by rhamnose, IPTG, or lactose.

26. (New) The method according to claim 22, wherein N-carbamoyl-L-methionine serve as substrates for the L-N-carbamoylase.

27. (New) The method according to claim 22, wherein the isolated polynucleotide is the *hyuC* gene of *Arthrobacter aurescens*.